

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

EFS

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/571,989A
Source: IFWO
Date Processed by STIC: 3/28/07

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/571,989A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence. (see item 11 below)

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWO

RAW SEQUENCE LISTING

DATE: 03/28/2007

PATENT APPLICATION: US/10/571,989A

TIME: 14:54:06

Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt

Output Set: N:\CRF4\03282007\J571989A.raw

3 <110> APPLICANT: Cleveland State University
 5 <120> TITLE OF INVENTION: EXOSITE-DIRECTED THROMBIN INHIBITORS
 7 <130> FILE REFERENCE: CLEV200023
 9 <140> CURRENT APPLICATION NUMBER: US 10/571,989A
 C--> 10 <141> CURRENT FILING DATE: 2006-03-13
 12 <150> PRIOR APPLICATION NUMBER: PCT/US04/021487
 13 <151> PRIOR FILING DATE: 2004-07-01
 15 <150> PRIOR APPLICATION NUMBER: US 60/502186
 16 <151> PRIOR FILING DATE: 2003-09-12
 18 <160> NUMBER OF SEQ ID NOS: 25
 20 <170> SOFTWARE: PatentIn version 3.2
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 30
 24 <212> TYPE: PRT
 25 <213> ORGANISM: Homo sapiens
 27 <400> SEQUENCE: 1
 29 Lys Met His Asp Arg Leu Glu Pro Gln Asp Glu Glu Ser Asp Ala Asp
 30 1 5 10 15
 33 Tyr Asp Tyr Gln Asn Arg Leu Ala Ala Ala Leu Gly Ile Arg
 34 20 25 30
 37 <210> SEQ ID NO: 2
 38 <211> LENGTH: 10
 39 <212> TYPE: PRT
 40 <213> ORGANISM: Homo sapiens
 42 <400> SEQUENCE: 2
 44 Lys Met His Asp Arg Leu Glu Pro Glu Asp
 45 1 5 10
 48 <210> SEQ ID NO: 3
 49 <211> LENGTH: 10
 50 <212> TYPE: PRT
 51 <213> ORGANISM: Homo sapiens
 53 <400> SEQUENCE: 3
 55 Leu Glu Pro Glu Asp Glu Glu Ser Asp Ala
 56 1 5 10
 59 <210> SEQ ID NO: 4
 60 <211> LENGTH: 10
 61 <212> TYPE: PRT
 62 <213> ORGANISM: Homo sapiens
 64 <400> SEQUENCE: 4
 66 Glu Glu Ser Asp Ala Asp Tyr Asp Tyr Gln
 67 1 5 10
 70 <210> SEQ ID NO: 5
 71 <211> LENGTH: 10

see pp 2-4
 Does Not Comply
 Corrected Diskette Needed

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Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt

Output Set: N:\CRF4\03282007\J571989A.raw

72 <212> TYPE: PRT
 73 <213> ORGANISM: Homo sapiens
 75 <400> SEQUENCE: 5
 77 Asp Tyr Asp Tyr Gln Asn Arg Leu Ala Ala
 78 1 5 10
 81 <210> SEQ ID NO: 6
 82 <211> LENGTH: 10
 83 <212> TYPE: PRT
 84 <213> ORGANISM: Homo sapiens
 86 <400> SEQUENCE: 6
 88 Asn Arg Leu Ala Ala Ala Leu Gly Ile Arg
 89 1 5 10
 92 <210> SEQ ID NO: 7
 93 <211> LENGTH: 5
 94 <212> TYPE: PRT
 95 <213> ORGANISM: Artificial
 97 <220> FEATURE:
 98 <223> OTHER INFORMATION: chemically synthesized
 100 <400> SEQUENCE: 7
 102 Asp Tyr Asp Tyr Gln
 103 1 5
 106 <210> SEQ ID NO: 8
 107 <211> LENGTH: 5
 108 <212> TYPE: PRT
 109 <213> ORGANISM: Artificial
 111 <220> FEATURE:
 112 <223> OTHER INFORMATION: TYR AT SEQUENCE LOCATION 4 IS SULFONATED (-SO3-2)
 114 <400> SEQUENCE: 8
 116 Asp Tyr Asp Tyr Gln
 117 1 5
 120 <210> SEQ ID NO: 9
 121 <211> LENGTH: 5
 122 <212> TYPE: PRT
 123 <213> ORGANISM: Artificial
 125 <220> FEATURE:
 126 <223> OTHER INFORMATION: TYR AT SEQUENCE LOCATIONS 2 AND 4 ARE SULFONATED (-SO3-2)
 128 <400> SEQUENCE: 9
 130 Asp Tyr Asp Tyr Gln
 131 1 5
 134 <210> SEQ ID NO: 10
 135 <211> LENGTH: 4
 136 <212> TYPE: PRT
 137 <213> ORGANISM: Homo sapiens
 139 <400> SEQUENCE: 10
 141 Asp Tyr Asp Tyr
 142 1
 145 <210> SEQ ID NO: 11
 146 <211> LENGTH: 5
 147 <212> TYPE: PRT

this may explain Tyr, but it does not explain
↓
Artificial Sequence
(see item 11 on Error Summary sheet)
same error

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Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt

Output Set: N:\CRF4\03282007\J571989A.raw

148 <213> ORGANISM: Homo sapiens
 150 <400> SEQUENCE: 11
 152 Asp Tyr Asp Tyr Gln
 153 1 5
 156 <210> SEQ ID NO: 12
 157 <211> LENGTH: 4
 158 <212> TYPE: PRT
 159 <213> ORGANISM: Artificial *same*
 161 <220> FEATURE:
 162 <223> OTHER INFORMATION: TYR AT SEQUENCE LOCATION 2 IS SULFONATED (-SO3-2)
 164 <400> SEQUENCE: 12
 166 Asp Tyr Asp Tyr
 167 1
 170 <210> SEQ ID NO: 13
 171 <211> LENGTH: 4
 172 <212> TYPE: PRT
 173 <213> ORGANISM: Artificial *same*
 175 <220> FEATURE:
 176 <223> OTHER INFORMATION: TYR AT SEQUENCE LOCATION 4 IS SULFONATED (-SO3-2)
 178 <400> SEQUENCE: 13
 180 Asp Tyr Asp Tyr
 181 1
 184 <210> SEQ ID NO: 14
 185 <211> LENGTH: 4
 186 <212> TYPE: PRT
 187 <213> ORGANISM: Artificial *same*
 189 <220> FEATURE:
 190 <223> OTHER INFORMATION: TYR AT SEQUENCE LOCATIONS 2 AND 4 ARE SULFONATED (-SO3-2)
 192 <400> SEQUENCE: 14
 194 Asp Tyr Asp Tyr
 195 1
 198 <210> SEQ ID NO: 15
 199 <211> LENGTH: 23
 200 <212> TYPE: DNA
 201 <213> ORGANISM: Homo sapiens
 203 <400> SEQUENCE: 15
 204 gagtgatgct aagtttgatt acc 23
 207 <210> SEQ ID NO: 16
 208 <211> LENGTH: 23
 209 <212> TYPE: DNA
 210 <213> ORGANISM: Homo sapiens
 212 <400> SEQUENCE: 16
 213 ggtaaatcaaa cttagcatca ctc 23
 216 <210> SEQ ID NO: 17
 217 <211> LENGTH: 18
 218 <212> TYPE: DNA
 219 <213> ORGANISM: Homo sapiens
 221 <400> SEQUENCE: 17
 222 catggagtga ccttctcg 18

RAW SEQUENCE LISTING

DATE: 03/28/2007

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Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt

Output Set: N:\CRF4\03282007\J571989A.raw

225 <210> SEQ ID NO: 18
 226 <211> LENGTH: 15
 227 <212> TYPE: DNA
 228 <213> ORGANISM: Homo sapiens
 230 <400> SEQUENCE: 18
 231 tcatccagga gaacc 15
 234 <210> SEQ ID NO: 19
 235 <211> LENGTH: 28
 236 <212> TYPE: DNA
 237 <213> ORGANISM: Homo sapiens
 239 <400> SEQUENCE: 19
 240 gctaagttta agttccagaa cagactgg 28
 243 <210> SEQ ID NO: 20
 244 <211> LENGTH: 28
 245 <212> TYPE: DNA
 246 <213> ORGANISM: Homo sapiens
 248 <400> SEQUENCE: 20
 249 ccagtctggt ctggaactta aacttagc 28
 252 <210> SEQ ID NO: 21
 253 <211> LENGTH: 13
 254 <212> TYPE: PRT
 255 <213> ORGANISM: Artificial
 257 <220> FEATURE:
 258 <223> OTHER INFORMATION: NO COMMENT
 260 <400> SEQUENCE: 21
 262 Asp Tyr Gln Asn Arg Leu Ala Ala Ala Leu Gly Ile Arg
 263 1 5 10
 266 <210> SEQ ID NO: 22
 267 <211> LENGTH: 15
 268 <212> TYPE: PRT
 269 <213> ORGANISM: Artificial
 271 <220> FEATURE:
 272 <223> OTHER INFORMATION: NO COMMENT
 274 <400> SEQUENCE: 22
 276 Pro Val Ile Pro Ala Asn Met Asp Lys Lys Tyr Arg Ser Gln His
 277 1 5 10 15
 280 <210> SEQ ID NO: 23
 281 <211> LENGTH: 42
 282 <212> TYPE: PRT
 283 <213> ORGANISM: Homo sapiens
 285 <400> SEQUENCE: 23
 287 Asn Leu Lys Lys Ile Thr Arg Glu Gln Arg Arg His Met Lys Arg Trp
 288 1 5 10 15
 291 Glu Tyr Phe Ile Ala Ala Glu Glu Val Ile Trp Asp Tyr Ala Pro Val
 292 20 25 30
 295 Ile Pro Ala Asn Met Asp Lys Lys Tyr Arg
 296 35 40
 299 <210> SEQ ID NO: 24
 300 <211> LENGTH: 9

*Invalid. Artificial
 Sequence must be explained
 (see item 11 on Error
 Summary sheet)*

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Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt

Output Set: N:\CRF4\03282007\J571989A.raw

301 <212> TYPE: PRT
302 <213> ORGANISM: Homo sapiens
304 <400> SEQUENCE: 24
306 Glu Tyr Phe Ile Ala Ala Glu Glu Val
307 1 5
310 <210> SEQ ID NO: 25
311 <211> LENGTH: 5
312 <212> TYPE: PRT
313 <213> ORGANISM: Homo sapiens
315 <400> SEQUENCE: 25
317 Glu Tyr Phe Ile Ala
318 1 5

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/571,989A

DATE: 03/28/2007
TIME: 14:54:07

Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt
Output Set: N:\CRF4\03282007\J571989A.raw

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#: 7, 8, 9, 12, 13, 14, 21, 22

VERIFICATION SUMMARY

DATE: 03/28/2007

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Input Set : N:\efs\03_28_07\10571989A_efs\clev200023ST25.txt

Output Set: N:\CRF4\03282007\J571989A.raw

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date